Page 1 of 6//

1644

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/503,421

DATE: 07/05/2000 TIME: 11:27:43

Input Set : A:\1970287.app

Output Set: N:\CRF3\07052000\1503421.raw

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3 <110> APPLICANT: SCHWAEBLE, Wilhelm
                University of Leicester, The
       6 <120> TITLE OF INVENTION: Clq and Collectin Receptor
       8 <130> FILE REFERENCE: M97/0287/PCT
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/503,421
  -> 11 <141> CURRENT FILING DATE: 2000-02-14-
      13 <160> NUMBER OF SEQ ID NOS: 7
      15 <170> SOFTWARE: PatentIn Ver. 2.0
      17 <210> SEQ ID NO: 1
      18 <211> LENGTH: 122
      19 <212> TYPE: PRT
      20 <213> ORGANISM: Homo sapiens
      22 <400> SEQUENCE: 1
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      26 Pro Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu Ser 27 \phantom{\bigg|}20\phantom{\bigg|}25\phantom{\bigg|}30\phantom{\bigg|}
      29 Gly Ser Leu Glu Asp Asp Trp Asp Phe Leu Pro Pro Lys Lys Ile Lys 30 \phantom{-}35\phantom{+}40\phantom{+}45\phantom{+}
      32 Asp Pro Asp Ala Ser Lys Pro Glu Asp Trp Asp Glu Arg Ala Lys Ile 33 \phantom{1}50\phantom{0} 55 \phantom{1}60\phantom{0}
      35 Asp Asp Pro Thr Asp Ser Lys Pro Glu Asp Trp Asp Lys Pro Glu His
36 65 70 75 80
      38 Ile Pro Asp Pro Asp Ala Lys Lys Pro Glu Asp Trp Asp Glu Glu Met
39 90 95
      41 Asp Gly Glu Trp Glu Pro Pro Val Ile Gln Asn Pro Glu Tyr Lys Gly 42 100 105 110
      44 Glu Trp Lys Pro Arg Gln Ile Asp Asn Pro
45 115 120
      48 <210> SEQ ID NO: 2
      49 <211> LENGTH: 122
      50 <212> TYPE: PRT
      51 <213> ORGANISM: Mus musculus
      53 <400> SEQUENCE: 2
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     57 Gln Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu Ser 58 20 25 30
      60 Gly Ser Leu Glu Asp Asp Gly Asp Phe Leu Pro Pro Lys Lys Ile Lys 61 \phantom{+}35\phantom{+}40\phantom{+}45\phantom{+}
      63 Asp Pro Asp Ala Ala Lys Pro Glu Asp Trp Asp Glu Arg Ala Lys Ile 64~50~55~60
      66 Asp Asp Pro Thr Asp Ser Lys Pro Glu Asp Trp Asp Lys Pro Glu His
67 65 70 75 80
      69 Ile Pro Asp Pro Asp Ala Lys Lys Pro Glu Asp Trp Asp Glu Glu Met
      72 Asp Gly Glu Trp Glu Pro Pro Val Ile Gln Asn Pro Glu Tyr Lys Gly
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ENTERED

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                 Input Set : A:\1970287.app
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75 Glu Trp Lys Pro Arg Gln Ile Asp Asn Pro
76 115 120
79 <210> SEQ ID NO: 3
80 <211> LENGTH: 122
81 <212> TYPE: PRT
82 <213> ORGANISM: Rattus norvegicus
84 <400> SEQUENCE: 3
85 Arg Cys Lys Asp Asp Glu Phe Thr His Leu Tyr Thr Leu Ile Val Arg 86 1 5 10 15
.88 Pro Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu Ser
89 20 25 30
91 Gly Ser Leu Glu Asp Asp Trp Asp Phe Leu Pro Pro Lys Lys Ile Lys
92 35 40 45
94 Asp Pro Asp Ala Ala Lys Pro Glu Asp Trp Asp Glu Arg Ala Lys Ile
95 50 60
97 Asp Asp Pro Thr Asp Ser Lys Pro Glu Asp Trp Asp Lys Pro Glu His
98 65 70 75 80
100 Tle Pro Asp Pro Asp Ala Lys Lys Pro Glu Asp Trp Asp Glu Glu Met 101 85 90 95
103 Asp Gly Glu Trp Glu Pro Pro Val Ile Gln Asn Pro Glu Tyr Lys Gly 104 \phantom{\bigg|}100\phantom{\bigg|} 105 \phantom{\bigg|}110\phantom{\bigg|}
106 Glu Trp Lys Pro Arg Gln Ile Asp Asn Pro
107 115
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111 <211> LENGTH: 366
112 <212> TYPE: DNA
113 <213> ORGANISM: Homo sapiens
115 <400> SEQUENCE: 4
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117 tatgaggtga agattgacaa cagccaggtg gagtccggct ccttggaaga cgattgggac 120
118 ttcctgccac ccaagaagat aaaggateet gatgetteaa aaceggaaga etgggatgag 180
119 egggecaaga tegatgatee cacagactee aageetgagg actgggacaa geecgageat 240
120 atccctgacc ctgatgctaa gaagcccgag gactgggatg aagagatgga cggagagtgg 300
121 gaaccccag tgattcagaa ccctgagtac aagggtgagt ggaagccccg gcagatcgac 360
122 aaccca
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125 <211> LENGTH: 366
126 <212> TYPE: DNA
127 <213> ORGANISM: Mus musculus
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130 cggtgtaagg atgatgaatt cacacaccta tacacactga ttgtgcggca agacaacacc 60
131 tatgaggtga aaattgacaa cagccaggtg gagtcaggct ccttggagga tgatggggac 120
132 tttctgccac ccaagaagat aaaggaccct gatgctgcca agccggaaga ctgggatgaa 180
133 cgagccaaga tcgatgaccc cacagattcc aagcctgagg actgggacaa gccagagcac 240
134 atccctgacc ctgatgctaa gaagcctgag gactgggatg aagagatgga tggagagtgg 300
135 gaaccaccag tgattcaaaa teetgaatac aagggegagt ggaaaccacg tcaaattgac 360
136 aaccca
138 <210> SEQ ID NO: 6
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RAW SEQUENCE LISTING

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140 <212> TYPE: DNA
141 <213> ORGANISM: Rattus norvegicus
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146 tttctgccgc ccaagaagat taaggatect gacgetgcca agccagaaga etgggatgaa 180
147 cgagccaaga ttgatgaccc cacagattcc aagcctgagg actgggacaa gccagagcac 240
148 atocotgaco otgatgotaa gaagootgag gaotgggacg aagagatgga tggagagtgg 300
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150 aaccca
152 <210> SEQ ID NO: 7
153 <211> LENGTH: 417
154 <212> TYPE: PRT
155 <213> ORGANISM: Homo sapiens
157 <400> SEQUENCE: 7
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161 Ala Glu Pro Ala Val Tyr Phe Lys Glu Gln Phe Leu Asp Gly Asp Gly 162 20 25 30
164 Trp Thr Pro Arg Trp Ile Glu Ser Lys His Lys Ser Asp Phe Gly Lys 165 \phantom{\bigg|} 35 \phantom{\bigg|} 40 \phantom{\bigg|} 45
167 Phe Val Leu Ser Ser Gly Lys Phe Tyr Gly Asp Glu Glu Lys Asp Lys 168 \phantom{\bigg|}50\phantom{\bigg|}
170 Gly Leu Gln Thr Ser Gln Asp Ala Arg Phe Tyr Ala Leu Ser Ala Ser
171 65 70 75 80
173 Phe Glu Pro Phe Ser Asn Lys Gly Gln Thr Leu Val Val Gln Phe Thr
174 85 90 95
176 Val Lys His Glu Gln Asn Ile Asp Cys Gly Gly Gly Tyr Val Lys Leu
177 100 105 110
179 Phe Pro Asn Ser Leu Asp Gln Thr Asp Met His Gly Asp Ser Glu Tyr
180 115 120 125
182 Asn Ile Met Phe Gly Pro Asp Ile Cys Gly Pro Gly Thr Lys Lys Val
183 130 135 140
185 His Val Ile Phe Asn Tyr Lys Gly Lys Asn Val Leu Ile Asn Lys Asp 186 145 150 150 155 160
188 Ile Arg Cys Lys Asp Asp Glu Phe Thr His Leu Tyr Thr Leu Ile Val
189 165 170 175

191 Arg Pro Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu
192 180 185 190
194 Ser Gly Ser Leu Glu Asp Asp Trp Asp Phe Leu Pro Pro Lys Lys Ile
195 195 200 205
197 Lys Asp Pro Asp Ala Ser Lys Pro Glu Asp Trp Asp Glu Arg Ala Lys
198 210 215 220
200 Ile Asp Asp Pro Thr Asp Ser Lys Pro Glu Asp Trp Asp Lys Pro Glu 201 225 230 235 240
203 His Ile Pro Asp Pro Asp Ala Lys Lys Pro Glu Asp Trp Asp Glu Glu 204 245 250 255
206 Met Asp Gly Glu Trp Glu Pro Pro Val Ile Gln Asn Pro Glu Tyr Lys
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207	260		265	270
209 Gly Glu 210	Trp Lys P 275	-	Ile Asp Asn Pro 280	Asp Tyr Lys Gly Thr 285
212 Trp Ile 213 290	His Pro G	lu Ile Asp 2 295		Ser Pro Asp Pro Ser 300
215 Ile Tyr 216 305	Ala Tyr A	sp Asn Phe 0 310	Gly Val Leu Gly 315	Leu Asp Leu Trp Gln 320
218 Val Lys 219	•	hr Ile Phe 2 25	Asp Asn Phe Leu 330	Ile Thr Asn Asp Glu 335
•		lu Phe Gly A	Asn Glu Thr Trp	Gly Val Thr Lys Ala
222	340		345	350
				350 Glu Gln Arg Leu Lys 365
224 Ala Glu 225	Lys Gln M 355	:	Lys Gln Asp Glu 360	Glu Gln Arg Leu Lys
224 Ala Glu 225 227 Glu Glu 228 370	Lys Gln M 355 Glu Glu A	sp Lys Lys 1 375	Lys Gln Asp Glu 360 Arg Lys Glu Glu	Glu Gln Arg Leu Lys 365 Glu Glu Ala Glu Asp
224 Ala Glu 225 227 Glu Glu 228 370 230 Lys Glu 231 385	Lys Gln M 355 Glu Glu A Asp Asp G	sp Lys Lys A 375 lu Asp Lys A 390	Lys Gln Asp Glu 360 Arg Lys Glu Glu Asp Glu Asp Glu 395	Glu Gln Arg Leu Lys 365 Glu Glu Ala Glu Asp 380 Glu Asp Glu Glu Asp

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/503,421

DATE: 07/05/2000 TIME: 11:27:44

Input Set : A:\1970287.app
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